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Sequence Listing could not be accepted.

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Reviewer: markspencer

Timestamp: Wed Jun 06 14:51:38 EDT 2007

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Reviewer Comments:

Applicant needs to remove all headers from each numeric identifier.

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Application No: 10567330

Version No: 1.0

Input Set:

Output Set:

Started: 2007-06-05 17:30:56.209

Finished: 2007-06-05 17:30:57.771

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 562 ms

Total Warnings: 20

Total Errors: 0

No. of SeqIDs Defined: 28

Actual SeqID Count: 28

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (27)
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**Error Description**

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<110> APPLICANT: Janatpour, Mary J.  
Reinhard, Christoph  
Garcia, Pablo  
<120> TITLE OF INVENTION: Trefoil Factor 3 (TFF3) as a Target for Anti-Cancer Therapy  
<130> FILE REFERENCE: CHIR0003-100 (19154.0006)

<140> CURRENT APPLICATION NUMBER:10567330  
<141> CURRENT FILING DATE:2007-06-05  
<150> PRIOR APPLICATION NUMBER: US/10/567,330  
<151> PRIOR FILING DATE: 2006-02-06  
<150> PRIOR APPLICATION NUMBER: US 60/493,173  
<151> PRIOR FILING DATE: 2003-08-07  
<150> PRIOR APPLICATION NUMBER: US 60/498,438  
<151> PRIOR FILING DATE: 2003-08-28  
<160> NUMBER OF SEQ ID NOS: 28  
<170> SOFTWARE: PatentIn version 3.2

<210> SEQ ID NO 1  
<211> LENGTH: 74  
<212> TYPE: PRT  
<213> ORGANISM: Homo sapiens  
<400> SEQUENCE: 1

Met	Leu	Gly	Leu	Val	Leu	Ala	Leu	Leu	Ser	Ser	Ser	Ser	Ala	Glu	Glu
1				5					10					15	
Tyr	Val	Gly	Leu	Ser	Ala	Asn	Gln	Cys	Ala	Val	Pro	Ala	Lys	Asp	Arg
			20					25					30		
Val	Asp	Cys	Gly	Tyr	Pro	His	Val	Thr	Pro	Lys	Glu	Cys	Asn	Asn	Arg
			35				40					45			
Gly	Cys	Cys	Phe	Asp	Ser	Arg	Ile	Pro	Gly	Val	Pro	Trp	Cys	Phe	Lys
	50					55				60					
Pro	Leu	Thr	Arg	Lys	Thr	Glu	Cys	Thr	Phe						
65					70										

<210> SEQ ID NO 2  
<211> LENGTH: 73  
<212> TYPE: PRT  
<213> ORGANISM: Homo sapiens  
<400> SEQUENCE: 2

Met	Leu	Gly	Leu	Val	Leu	Ala	Leu	Leu	Ser	Ser	Ser	Ser	Ala	Glu	Glu
1				5					10					15	
Tyr	Val	Gly	Leu	Ser	Ala	Asn	Gln	Cys	Ala	Val	Pro	Ala	Lys	Asp	Arg
			20					25					30		
Val	Asp	Cys	Gly	Tyr	Pro	His	Val	Thr	Pro	Lys	Glu	Cys	Asn	Asn	Arg
			35				40					45			
Gly	Cys	Cys	Phe	Asp	Ser	Arg	Ile	Pro	Gly	Val	Pro	Trp	Cys	Phe	Lys
	50					55				60					
Pro	Leu	Gln	Glu	Ala	Glu	Cys	Thr	Phe							
65					70										

<210> SEQ ID NO 3  
<211> LENGTH: 80  
<212> TYPE: PRT  
<213> ORGANISM: Homo sapiens  
<400> SEQUENCE: 3

Met	Ala	Ala	Arg	Ala	Leu	Cys	Met	Leu	Gly	Leu	Val	Leu	Ala	Leu	Leu
1				5					10					15	
Ser	Ser	Ser	Ser	Ala	Glu	Glu	Tyr	Val	Gly	Leu	Ser	Ala	Arg	Gly	Cys

	20		25		30										
Ala	Val	Pro	Ala	Lys	Asp	Arg	Val	Asp	Cys	Gly	Tyr	Pro	His	Val	Thr
	35		40		45										
Pro	Lys	Glu	Cys	Asn	Asn	Arg	Gly	Cys	Cys	Phe	Asp	Ser	Arg	Ile	Pro
	50		55		60										
Gly	Val	Pro	Trp	Cys	Phe	Lys	Pro	Leu	Gln	Glu	Ala	Glu	Cys	Thr	Phe
65			70					75						80	

<210> SEQ ID NO 4  
 <211> LENGTH: 130  
 <212> TYPE: PRT  
 <213> ORGANISM: Homo sapiens  
 <400> SEQUENCE: 4

Met	Gln	Glu	Arg	Thr	Gly	Ala	Ala	Thr	Ala	Arg	Arg	Glu	Ser	Leu	Pro
1				5				10						15	
Gln	Ala	Asn	Asn	Pro	Glu	Gln	Leu	Cys	Lys	Gln	Arg	Cys	Ile	Asn	Glu
			20					25					30		
Ala	Ser	Trp	Thr	Met	Lys	Arg	Val	Leu	Ser	Cys	Val	Pro	Glu	Pro	Thr
	35						40						45		
Val	Val	Met	Ala	Ala	Arg	Ala	Leu	Cys	Met	Leu	Gly	Leu	Val	Leu	Ala
	50					55					60				
Leu	Leu	Ser	Ser	Ser	Ser	Ala	Glu	Glu	Tyr	Val	Gly	Leu	Ser	Ala	Asn
65					70				75					80	
Gln	Cys	Ala	Val	Pro	Ala	Lys	Asp	Arg	Val	Asp	Cys	Gly	Tyr	Pro	His
			85					90						95	
Val	Thr	Pro	Lys	Glu	Cys	Asn	Asn	Arg	Gly	Cys	Cys	Phe	Asp	Ser	Arg
		100						105						110	
Ile	Pro	Gly	Val	Pro	Trp	Cys	Phe	Lys	Pro	Leu	Gln	Glu	Ala	Glu	Cys
		115						120						125	
Thr	Phe														
	130														

<210> SEQ ID NO 5  
 <211> LENGTH: 398  
 <212> TYPE: DNA  
 <213> ORGANISM: Homo sapiens  
 <400> SEQUENCE: 5

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gtctgcaaac	cagtgtgccg	tgccggccaa	ggacagggtg	gactgcggtc	acccccatgt	120
cacccccaa	gagtgcaca	accggggctg	ctgctttgac	tccaggatcc	ctggagtgcc	180
ttggtgtttc	aagccctga	ctaggaagac	agaatgcacc	ttctgaggca	cctccagctg	240
cccctgggat	gcaggtctg	cacccttgcc	cggtctgat	tgtgtccagg	cactgttcat	300
ctcagttttt	ctgtcccttt	gctcccggca	agctttctgc	tgaaagttca	tatctggagc	360
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 <213> ORGANISM: Homo sapiens  
 <400> SEQUENCE: 6

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caagcaaaca	atccagagca	gctgtgcaaa	caacgggtgca	taaatgaggc	ctcctggacc	180
atgaagcgag	tcctgagctg	cgtcccggag	cccacgggtg	tcatggctgc	cagagcgctc	240
tgcattgctg	ggctggctct	ggccttgctg	tcctccagct	ctgctgagga	gtacgtgggc	300
ctgtctgcaa	accagtgtgc	cgtgccagcc	aaggacaggg	tggactgcgg	ctacccccat	360
gtcaccccc	aggagtgc	caaccggggc	tgtctgcttg	actccaggat	ccctggagtg	420

ccttggtggt	tcaagccct	gcaggaagca	gaatgcacct	tctgaggcac	ctccagctgc	480
cccgcgccg	gggatgcgag	gctcggagca	cccttgcccg	gctgtgattg	ctgccaggca	540
ctgttcatct	cagcttttct	gtccctttgc	tcccggaag	cgcttctgct	gaaagtcat	600
atctggagcc	tgatgtctta	acgaataaag	gtcccatgct	ccaccgagg	acagttcttc	660
gtgcctgaaa	aaaaaaaaaa	aaaaa				685

<210> SEQ ID NO 7  
 <211> LENGTH: 491  
 <212> TYPE: DNA  
 <213> ORGANISM: Homo sapiens  
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ctggggctgg	tcctggcctt	gctgtcctcc	agctctgctg	aggagtacgt	gggcctgtct	120
gcaaaccagt	gtgccgtgcc	agccaaggac	aggggtggact	gcggctaccc	ccatgtcacc	180
cccaaggagt	gcaacaaccg	gggctgctgc	tttgactcca	ggatccctgg	agtgccttgg	240
tgtttcaagc	ccctgcagga	agcagaatgc	accttctgag	gcacctccag	ctgcccccg	300
ccgggggatg	cgaggctcgg	agcacccttg	ccgggtgtg	attgctgcc	ggcactgttc	360
atctcagctt	ttctgtccct	ttgtcccg	caagcgcttc	tgctgaaagt	tcatatctgg	420
agcctgatgt	cttaacgaat	aaagggtcca	tgctccacc	taaaaaaaaa	aaaaaaaaaa	480
aaaaaaaaaa	a					491

<210> SEQ ID NO 8  
 <211> LENGTH: 432  
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aacagtgggg	gctgaactga	cctctccct	ttgggagaga	aaaactgtct	gggagcttga	120
caaaggcatg	caggagagaa	caggagcagc	cacagccagg	agggagagcc	ttccccagc	180
aaacaatcca	gagcagctgt	gcaaacaacg	gtgcataaat	gaggcctcct	ggaccatgaa	240
gcgagtcttg	agctgcgtcc	cggagcccac	ggtggatcatg	gctgccagag	cgctctgcat	300
gctggggctg	gtcctggcct	tgctgtcctc	cagctctgct	gaggagtacg	tgggcctgtc	360
tgcaaaccag	tgtgccgtgc	cagccaagga	caggggtggac	tgcggtacc	cccatgtcac	420
ccccaaggag	tg					432

<210> SEQ ID NO 9  
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 <220> FEATURE:  
 <223> OTHER INFORMATION: TFF3 antisense oligonucleotide  
 <400> SEQUENCE: 9

tccttggtg	gcacgcaca	ct	22
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<210> SEQ ID NO 10  
 <211> LENGTH: 23  
 <212> TYPE: DNA  
 <213> ORGANISM: Artificial sequence  
 <220> FEATURE:  
 <223> OTHER INFORMATION: TFF3 antisense oligonucleotide  
 <400> SEQUENCE: 10

cgggagcaaa	gggacagaaa	agc	23
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<210> SEQ ID NO 11  
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 <212> TYPE: DNA  
 <213> ORGANISM: Artificial sequence

<220> FEATURE:  
 <223> OTHER INFORMATION: TFF3 antisense oligonucleotide  
 <400> SEQUENCE: 11  
 gaagaactgt cctcgggtgg agc 23

<210> SEQ ID NO 12  
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 tcagaaagtc tcaggcacga agaac 25

<210> SEQ ID NO 13  
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 <213> ORGANISM: Artificial sequence  
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 gcagcagaaa taaagcacia cctca 25

<210> SEQ ID NO 14  
 <211> LENGTH: 25  
 <212> TYPE: DNA  
 <213> ORGANISM: Artificial sequence  
 <220> FEATURE:  
 <223> OTHER INFORMATION: TFF3 antisense oligonucleotide  
 <400> SEQUENCE: 14  
 aacagtagcg agagtgggtg tgaaa 25

<210> SEQ ID NO 15  
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 <213> ORGANISM: Artificial sequence  
 <220> FEATURE:  
 <223> OTHER INFORMATION: TFF3 antisense oligonucleotide  
 <400> SEQUENCE: 15  
 cggcacggca cactggtttg ca 22

<210> SEQ ID NO 16  
 <211> LENGTH: 25  
 <212> TYPE: DNA  
 <213> ORGANISM: Artificial sequence  
 <220> FEATURE:  
 <223> OTHER INFORMATION: TFF3 antisense oligonucleotide  
 <400> SEQUENCE: 16  
 ggtgcattct gtcttcctag tcagg 25

<210> SEQ ID NO 17  
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 <212> TYPE: DNA  
 <213> ORGANISM: Artificial sequence  
 <220> FEATURE:  
 <223> OTHER INFORMATION: TFF3 antisense oligonucleotide  
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ggctccagat atgaactttc agcag

25

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<212> TYPE: DNA  
<213> ORGANISM: Artificial sequence  
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ggtggagcat gggaccttta ttcgt

25

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<212> TYPE: DNA  
<213> ORGANISM: Artificial sequence  
<220> FEATURE:  
<223> OTHER INFORMATION: TFF3 antisense oligonucleotide  
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tggcacggca cactggtttg ca

22

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Ala Val Pro Ala Lys Asp Arg Val  
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Val Pro Ala Lys Asp Arg Val Asp  
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<212> TYPE: PRT  
<213> ORGANISM: Artificial sequence  
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<223> OTHER INFORMATION: chemically synthesized peptide  
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Ala Val Pro Ala Lys Asp Arg Val Asp  
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<212> TYPE: PRT  
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<220> FEATURE:  
<223> OTHER INFORMATION: chemically synthesized peptide  
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Gly Tyr Pro His Val Thr Pro Lys  
1 5

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Tyr Pro His Val Thr Pro Lys Glu  
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<213> ORGANISM: Artificial sequence  
<220> FEATURE:  
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Gly Tyr Pro His Val Thr Pro Lys Glu  
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Phe Lys Pro Leu Gln Glu Ala Glu  
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<220> FEATURE:  
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Lys Pro Leu Gln Glu Ala Glu Cys  
1 5

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<220> FEATURE:  
<223> OTHER INFORMATION: chemically synthesized peptide  
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Phe Lys Pro Leu Gln Glu Ala Glu Cys  
1 5